

0400

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/780,668

Source: 0//E

Date Processed by STIC: 3/7/2,00/

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE
APPLICANT, WITH A NOTICE TO COMPLY or,

2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A

FOR CRESUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help://patin21help@uspto.gov/or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help://patin3help@uspto.gov/or phone 703-306-4119 (R. Wax)

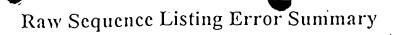
TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

## **Checker Version 3.0**

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker



SERIAL NUMBER: 09/

## ERROR DETECTED SUGGESTED CORRECTION

1	Wrapped Nucleics	The number/text at the end of each line "wrapped" down to the next line.
	•	This may occur if your file was retrieved in a word processor after creating it.  Please adjust your right margin to .3, as this will prevent "wrapping".
2	Wrapped Aminos	The amino acid number/text at the end of each line "wrapped" down to the next line.
		This may occur if your file was retrieved in a word processor after creating it.  Please adjust your right margin to .3, as this will prevent "wrapping".
3	Incorrect Line Length	The rules require that a line not exceed 72 characters in length. This includes spaces.
4	Misaligned Amino Acid Numbering	The numbering under each 5th amino acid is misaligned. This may be caused by the use of tabs between the numbering. It is recommended to delete any tabs and use spacing between the numbers.
5	Non-ASCII	This file was not saved in ASCII (DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text so that it can be processed.
6	Variable Length	Sequence(s) contain n's or Xaa's which represented more than one residue.  As per the rules, each n or Xaa can only represent a single residue.
		Please present the maximum number of each residue having variable length and indicate in the (ix) feature section that some may be missing.
<sup>7</sup> ——	Patentin ver. 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequence(s) Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies primarily to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
8	Skipped Sequences (OLD RULES)	Sequence(s) missing. If intentional, please use the following format for each skipped sequence:  (2) INFORMATION FOR SEQ ID NO:X:  (i) SEQUENCE CHARACTERISTICS:(Do not insert any headings under "SEQUENCE CHARACTERISTICS")  (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: This sequence is intentionally skipped
	•	Please also adjust the "(iii) NUMBER OF SEQUENCES:" response to include the skipped sequence(s).
9	Skipped Sequences (NEW RULES)	Sequence(s) missing. If intentional, please use the following format for each skipped sequence.  <210> sequence id number  <400> sequence id number  000
o	Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing.  Use of <220> to <223> is MANDATORY if n's or Xaa's are present.  In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
1	Use of <213>Organism (NEW RULES)	Sequence(s) are missing this mandatory field or its response.
· —	Use of <220>Feature (NEW RULES)	Sequence(s) are missing the <220>Feature and associated headings.  Use of <220> to <223> is MANDATORY if <213>ORGANISM is "Artificial" or "Unknown"  Please explain source of genetic material in <220> to <223> section.
3	Patentin ver. 2.0 "bug"	(See "Federal Register," 6/01/98, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of new Rules)  Please do not use "Copy to Disk" function of Patentin version 2.0. This causes a corrupted  file; resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing).

Instead, please use "File Manager" or any other means to copy file to floppy disk.

OIPE

```
Does Not Comply
                                                                                    Corrected Diskette Needed
                      Input Set : A:\LEX011.ST25.txt
                      Output Set: N:\CRF3\03072001\I780668.raw
      3 <110> APPLICANT: Gillies, Stephen
              Burger, Christa
              Lo, Kin-Ming
      7 <120> TITLE OF INVENTION: Enhancing the Circulating Half-Life of Antibody-Based Fusion Proteins
      9 <130> FILE REFERENCE: LEX-011
C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/780,668
C--> 11 <141> CURRENT FILING DATE: 2001-02-09
     11 <150> PRIOR APPLICATION NUMBER: US 60/181,768
     12 <151> PRIOR FILING DATE: 2000-02-11
     14 <160> NUMBER OF SEQ ID NOS: 35
     16 <170> SOFTWARE: PatentIn version 3.0
     18 <210> SEQ ID NO: 1
     19 <211> LENGTH: 7
     20 <212> TYPE: PRT
     21 <213> ORGANISM: Artificial Sequence
     23 <220> FEATURE:
     24 <223> OTHER INFORMATION: Ig-IL-2 junction sequence
     26 <400> SEQUENCE: 1
     28 Ser Pro Gly Lys Ala Pro Thr
     29 1
     31 <210> SEQ ID NO: 2
     32 <211> LENGTH: 4
     33 <212> TYPE: PRT
     34 <213> ORGANISM: Artificial Sequence
     36 <220> FEATURE:
     37 <223> OTHER INFORMATION: Ig C-terminal sequence
     39 <400> SEQUENCE: 2
     41 Ser Pro Gly Lys
     42 1
     44 <210> SEQ ID NO: 3
     45 <211> LENGTH: 12
     46 <212> TYPE: DNA
                                                      global resporse gerê source of

genetic.

genetic.

(see arrowed

portion of

for Jen 12 on

Evan Lumnaur

10 11
     47 <213> ORGANISM: Artificial Sequence
     49 <220> FEATURE:
     50 <223> OTHER INFORMATION: Synthetic sequence
     52 <400> SEQUENCE: 3
     53 tccccgggta aa
     56 <210> SEQ ID NO: 4
     57 <211> LENGTH: 42
     58 <212> TYPE: DNA
     59 <213> ORGANISM: Artificial Sequence
     61 <220> FEATURE:
     62 <223> OTHER INFORMATION: Synthetic sequence
     64 <400> SEQUENCE: 4
     65 ccgggtgcag cacctacttc aagttctaca aagaaaacac ag
     68 <210> SEQ ID NO: 5
     69 <211> LENGTH: 38
```

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/780,668

DATE: 03/07/2001

TIME: 11:13:52



RAW SEQUENCE LISTING DATE: 03/07/2001 PATENT APPLICATION: US/09/780,668 TIME: 11:13:52

Input Set : A:\LEX011.ST25.txt

Output Set: N:\CRF3\03072001\1780668.raw

70 <212> TYPE: DNA	
71 <213> ORGANISM: Artificial Sequence	
73 <220> FEATURE:	
74 <223> OTHER INFORMATION: Synthetic sequence	
76 <400> SEQUENCE: 5	
77 ctgtgttttc tttgtagaac ttgaagtagg tgctgcac	38
80 <210> SEQ ID NO: 6	
81 <211> LENGTH: 42	
82 <212> TYPE: DNA	
83 <213> ORGANISM: Artificial Sequence	
85 <220> FEATURE:	
86 <223> OTHER INFORMATION: Synthetic sequence	
88 <400> SEQUENCE: 6	
89 ccgggtaggg cgccaacttc aagttctaca aagaaaacac ag	42
92 <210> SEQ ID NO: 7	
93 <211> LENGTH: 38	
94 <212> TYPE: DNA	
95 <213> ORGANISM: Artificial Sequence	
97 <220> FEATURE:	
98 <223> OTHER INFORMATION: Synthetic sequence	
100 <400> SEQUENCE: 7	
101 ctgtgttttc tttgtagaac ttgaagttgg cgccctac	38
104 <210> SEQ ID NO: 8 `	
105 <211> LENGTH: 39	
106 <212> TYPE: DNA	
107 <213> ORGANISM: Artificial Sequence	
109 <220> FEATURE:	
110 <223> OTHER INFORMATION: Synthetic sequence	
112 <400> SEQUENCE: 8	
113 ccgggtgcac ctacttcaag ttctacaaag aaaacacag	39
116 <210> SEQ ID NO: 9	
117 <211> LENGTH: 35	
118 <212> TYPE: DNA	
119 <213> ORGANISM: Artificial Sequence	
121 <220> FEATURE:	
122 <223> OTHER INFORMATION: Synthetic sequence	
124 <400> SEQUENCE: 9	0.5
125 ctgtgttttc tttgtagaac ttgaagtagg tgcac	35
128 <210> SEQ ID NO: 10	
129 <211> LENGTH: 42	
130 <212> TYPE: DNA	
131 <213> ORGANISM: Artificial Sequence	
133 <220> FEATURE:	
134 <223> OTHER INFORMATION: Synthetic sequence	
136 <400> SEQUENCE: 10	42
137 cegggtgggg cecetactic aagttetaca aagaaaacac ag	42
140 <210> SEQ ID NO: 11	
141 <211> LENGTH: 38	
142 <212> TYPE: DNA	



RAW SEQUENCE LISTING DATE: 03/07/2001 PATENT APPLICATION: US/09/780,668 TIME: 11:13:52

Input Set : A:\LEX011.ST25.txt

Output Set: N:\CRF3\03072001\I780668.raw

143 <213> ORGANISM: Artificial Sequence 145 <220> FEATURE: 146 <223> OTHER INFORMATION: Synthetic sequence 148 <400> SEQUENCE: 11 38 -149 ctgtgttttc tttgtagaac ttgaagtagg ggccccac 152 <210> SEQ ID NO: 12 153 <211> LENGTH: 42 154 <212> TYPE: DNA 155 <213> ORGANISM: Artificial Sequence 157 <220> FEATURE: 158 <223> OTHER INFORMATION: Synthetic sequence 160 <400> SEQUENCE: 12 42 161 ccgggtctgg cgccaacttc aagttctaca aagaaaacac ag 164 <210> SEQ ID NO: 13 165 <211> LENGTH: 38 166 <212> TYPE: DNA 167 <213> ORGANISM: Artificial Sequence 169 <220> FEATURE: 170 <223> OTHER INFORMATION: Synthetic sequence 172 <400> SEQUENCE: 13 38 173 ctgtgttttc tttgtagaac ttgaagttgg cgccagac 176 <210> SEQ ID NO: 14 177 <211> LENGTH: 48 178 <212> TYPE: DNA 179 <213> ORGANISM: Artificial Sequence 181 <220> FEATURE: 182 <223> OTHER INFORMATION: Synthetic sequence 184 <400> SEQUENCE: 14 48 185 ccgggtgcag cagctgcccc aacttcaagt tctacaaaga aaacacag 188 <210> SEQ ID NO: 15 189 <211> LENGTH: 44 190 <212> TYPE: DNA 191 <213> ORGANISM: Artificial Sequence 193 <220> FEATURE: 194 <223> OTHER INFORMATION: Synthetic sequence 196 <400> SEQUENCE: 15 197 ctgtgttttc tttgtagaac ttgaagttgg ggcagctgct gcac 44 200 <210> SEQ ID NO: 16 201 <211> LENGTH: 42 202 <212> TYPE: DNA 203 <213> ORGANISM: Artificial Sequence 205 <220> FEATURE: 206 <223> OTHER INFORMATION: Synthetic sequence 208 <400> SEQUENCE: 16 42 209 ccgggttgcg caccaacttc aagttctaca aagaaaacac ag 212 <210> SEQ ID NO: 17 213 <211> LENGTH: 38 214 <212> TYPE: DNA

215 <213> ORGANISM: Artificial Sequence

DATE: 03/07/2001 RAW SEQUENCE LISTING PATENT APPLICATION: US/09/780,668 TIME: 11:13:52

Input Set : A:\LEX011.ST25.txt

Output Set: N:\CRF3\03072001\1780668.raw

```
217 <220> FEATURE:
     218 <223> OTHER INFORMATION: Synthetic sequence
     220 <400> SEQUENCE: 17
     221 ctgtgttttc tttgtagaac ttgaagttgg tgcgcaac
                                                                               38
     224 <210> SEQ ID NO: 18
     225 <211> LENGTH: 42
     226 <212> TYPE: DNA
     227 <213> ORGANISM: Artificial Sequence
     229 <220> FEATURE:
     230 <223> OTHER INFORMATION: Synthetic sequence
     232 <400> SEQUENCE: 18
                                                                               42
     233 cegggtgacg caccaacttc aagttctaca aagaaaacac ag
     236 <210> SEQ ID NO: 19
     237 <211> LENGTH: 38
     238 <212> TYPE: DNA
     239 <213> ORGANISM: Artificial Sequence
     241 <220> FEATURE:
     242 <223> OTHER INFORMATION: Synthetic sequence
     244 <400> SEQUENCE: 19
                                                                               38
     245 ctgtgttttc tttgtagaac ttgaagttgg tgcgtcac
     248 <210> SEQ ID NO: 20
     249 <211> LENGTH: 19
     250 <212> TYPE: DNA
     251 <213> ORGANISM: Artificial Sequence
     253 <220> FEATURE:
     254 <223> OTHER INFORMATION: Synthetic sequence
     256 <220> FEATURE:
     257 <221> NAME/KEY: CDS
     258 <222> LOCATION: (2)..(19)
     260 <400> SEQUENCE: 20
     261 c ccg gca tgc ggg ggt aaa
                                                                               19
     262
          Pro Ala Cys Gly Gly Lys
     263
          1
     266 <210> SEQ ID NO: 21
     267 <211> LENGTH: 6
                                                                     -) sel ten 2 on Evar
Summany Sheet
     268 <212> TYPE: PRT
     269 <213 ORGANISM: Artificial Sequence
W--> 271 (220) FEATURE:
271 $400> SEQUENCE: 21
     273 Pro Ala Cys Gly Gly Lys
     274 1.
     277 <210> SEQ ID NO: 22
     278 <211> LENGTH: 18
     279 <212> TYPE: DNA
     280 <213> ORGANISM: Artificial Sequence
     282 <220> FEATURE:
     283 <223> OTHER INFORMATION: Synthetic sequence
     285 <400> SEQUENCE: 22
```



RAW SEQUENCE LISTING DATE: 03/07/2001 PATENT APPLICATION: US/09/780,668 TIME: 11:13:52

Input Set : A:\LEX011.ST25.txt

Output Set: N:\CRF3\03072001\I780668.raw .

	gggttcagga tccggagg	18
289	<210> SEQ ID NO: 23	
290	<211> LENGTH: 18	
	<212> TYPE: DNA	
292	<213> ORGANISM: Artificial Sequence	
294	<220> FEATURE:	•
295	<223> OTHER INFORMATION: Synthetic sequence	
297	<400> SEQUENCE: 23	
298	cctccggatc ctgaaccc	18
301	<210> SEQ ID NO: 24	
302	<211> LENGTH: 9	
303	<212> TYPE: PRT	
304	<213> ORGANISM: Artificial Sequence	
306	<220> FEATURE:	
307	<223> OTHER INFORMATION: Synthetic sequence	
309	<400> SEQUENCE: 24	
311	Pro Gly Ser Gly Ser Gly Gly Lys	
312		
314	<210> SEQ ID NO: 25	
	<211> LENGTH: 33	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial Sequence	
	<220> FEATURE:	
	<223> OTHER INFORMATION: Synthetic sequence	
	<400> SEQUENCE: 25	
	gggttcaggc tctggatcag ggtccggatc cgg	33
	<210> SEQ ID NO: 26	
	<211> LENGTH: 33	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial Sequence	
	<220> FEATURE:	
	<223> OTHER INFORMATION: Synthetic sequence	
	<400> SEQUENCE: 26	
	ceggateegg accetgatee agageetgaa eee	33
	<210> SEQ ID NO: 27	•
	<211> LENGTH: 14	
	<211> DENGIN: 14 <212> TYPE: PRT	
	<213>ORGANISM: Artificial Sequence	
	<220> FEATURE:	
	<pre>&lt;220&gt; PERIORE. &lt;223&gt; OTHER INFORMATION: Synthetic sequence</pre>	
•	<400> SEQUENCE: 27	
	Pro Gly Ser Gly Ser Gly Ser Gly Ser Gly Gly Lys	
349		
	<210> SEQ ID NO: 28	
	<211> LENGTH: 25	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial Sequence	
	<220> FEATURE:	
33/	<223> OTHER INFORMATION: Synthetic sequence	

reach to the substitutions





VERIFICATION SUMMARY

PATENT APPLICATION: US/09/780,668

DATE: 03/07/2001 TIME: 11:13:53

Input Set : A:\LEX011.ST25.txt

Output Set: N:\CRF3\03072001\1780668.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application No L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:271 M:258 W: Mandatory Feature missing, <220> FEATURE:

L:271 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:



VERIFICATION SUMMARY

PATENT APPLICATION: US/09/780,668

DATE: 03/07/2001 TIME: 11:13:53

Input Set : A:\LEX011.ST25.txt

Output Set: N:\CRF3\03072001\I780668.raw .

 $L:11 \text{ M}:270 \text{ C}: \text{ Current Application Number differs, Replaced Current Application No} \\ L:11 \text{ M}:271 \text{ C}: \text{ Current Filing Date differs, Replaced Current Filing Date}$ 

L:271 M:258 W: Mandatory Feature missing, <220> FEATURE:

L:271 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: